CP DOMAIN OF THE IGF-1RECEPTOR \Im 궃 CP DOMAIN OF THE IGF-1RECEPTOR 2 = GAA TTC CCG GGG AGA AAG AGA B R K R CAA TTC AGA AAG AGA R K R AAG AGA K/A MUTATION AGA R GAA TTC CCG GGG E F P G LINKER LINKER CAA KINASE INACTIVE MUTANT LexA DNA BINDING DOMAIN Lexa DNA BINDING DOMAIN WILDTYPE

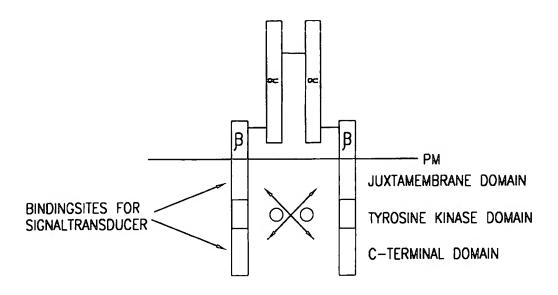
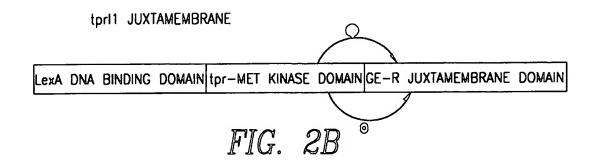
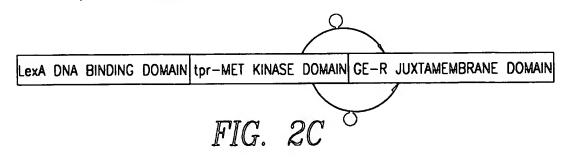


FIG. 2A



tprl1 C-TERMINAL



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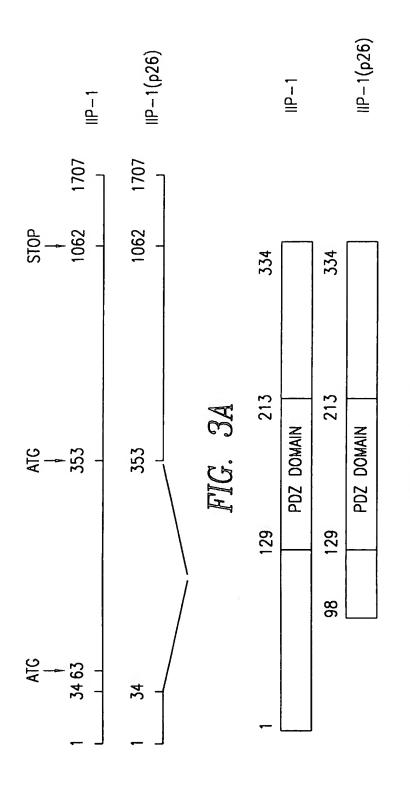
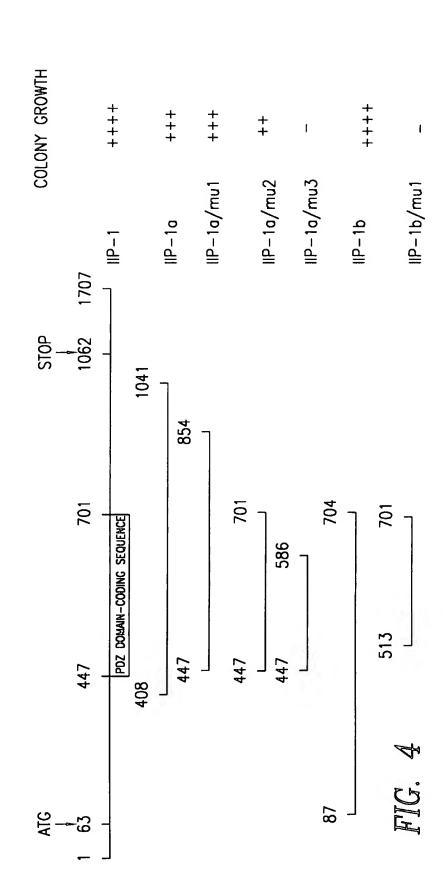


FIG. 3B



Hig. 5

- 1 MSRPRKRLAG TSGSBKGLSG KRTKTENSGE ALAKVEDSNP QKTSATKNCL
- 51 KNLSSHWLMK SEPESRLEKG VDVKFSIEDL KAQPKQTTEW DGVRNYQARN
- + FLRAMKLGEE AFFYHSNCKE PGIAGLMKIV KEAYPDHTÖF EKNNPHYDPS 101
- 151 SKEDNPKWSM VDVQFVRMMK RFIPLAELKS YHQAHKATGG PLKNMVLFTR
- 201 QRLSIQPLTQ EEFDFVLSLE EKEPS*
- ' putative N-glycosylation site
- putative Casein kinase II phosphorylation site
- putative N-myristoylation site
- putative Tyrosine kinase phosphorylation site
- ♦ putative Protein kinase C phosphorylation site
- **←→** putative NLS

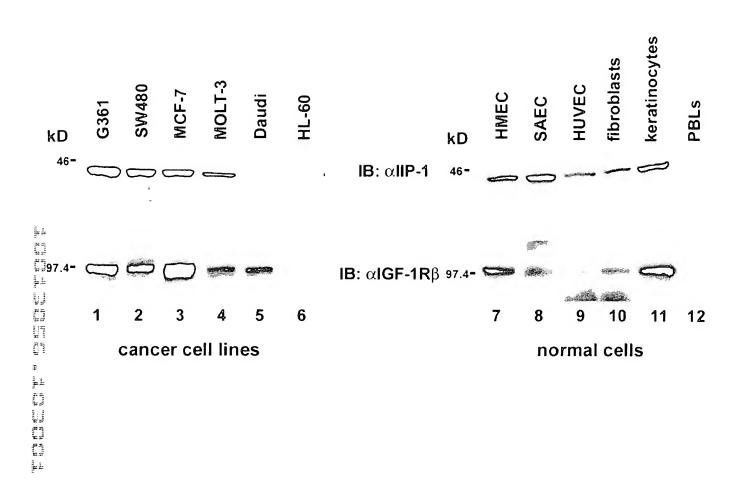






Fig. 7

